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Application No.: 10/785, 220

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

☒ 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.

☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).

☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).

☐ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."

☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).

☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).

☒ 7. Other: _See attachment._

Applicant Must Provide:

☒ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".

☒ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.

☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216
For CRF Submission Help, call (703) 308-4212
For PatentIn software help, call (703) 308-6856

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/785,220A
Source: FROM6
Date Processed by STIC: 9-6-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 10/785,200A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____ Wrapped Nucleics
Wrapped Aminos — The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .): this will prevent "wrapping."

2____ Invalid Line Length — The rules require that a line not exceed 72 characters in length. This includes white spaces.

3____ Misaligned Amino
Numbering — The numbering under each 5ᵗʰ amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4____ Non-ASCII — The submitted file was not saved in ASCII(DOS)text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5____ Variable Length — Sequence(s)____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<22]> section that some may be missing

6____ PatentIn 2.0
"bug" — A "bug" in PatentIn version 2.0 has caused the <220>-<22]> section to be missing from amino acid sequences(s)____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<22]> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<22]> sections for Artificial or Unknown sequences.

7____ Skipped Sequences
(OLD RULES) — Sequence(s)____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO X (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION SEQ ID NO X (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(xi) NUMBER OF SEQUENCES" response to include the skipped sequences.

8____ Skipped Sequences
(NEW RULES) — Sequence(s)____ missing. If intentional, please insert the following lines for each skipped sequence:
<210> sequence id number
<400> sequence id number
000

9____ Use of n's or Xaa's
(NEW RULES) — Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<22]> is MANDATORY if n's or Xaa's are present. In <220> to <22]> section, please explain location of n or Xaa, and which residue n or Xaa represents

10____ Invalid <22]>
Response — Per 1.823 of Sequence Rules, the only valid <22]> responses are Unknown, Artificial Sequence or scientific name (Genus/species) <220> <22]> section is required when <22]> response is Unknown... is Artificial Sequence

11____ Use of <220> — Use of <220> to <22]> is MANDATORY if <22]> Organism response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <22]> section. (See "Federal Register," 06/01/1998, Vol 63, No 104, pp.29631-32) (Sec. 1.823 of Sequence Rules)

12____ PatentIn 2.0
"bug" — Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____ Misuse of n/Xaa — "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

AMC - Biotechnology Systems Branch - 09/09/2003
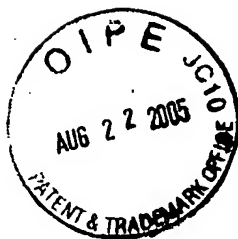
IFW16

RAW SEQUENCE LISTING   DATE: 04/06/2005
PATENT APPLICATION: US/10/785,220A   TIME: 14:13:57

Input Set : A:\39780-1216.TXT
Output Set: N:\CRF4\04062005\J785220A.raw

```
 4 <110> APPLICANT: Ashkenazi, Avi J.
 5           Fong, Sherman
 6           Goddard, Audrey
 7           Gurney, Austin L.
 8           Napier, Mary A.
 9           Tumas, Daniel
10           Wood, William I.
12 <120> TITLE OF INVENTION: COMPOUNDS, COMPOSITIONS AND METHODS FOR
13        THE TREATMENT OF DISEASES CHARACTERIZED BY A33- RELATED
14        ANTIGENS
16 <130> FILE REFERENCE: P1216R1C1D4
18 <140> CURRENT APPLICATION NUMBER: 10/785,220A
19 <141> CURRENT FILING DATE: 2004-02-24
21 <150> PRIOR APPLICATION NUMBER: US 09/254,465
22 <151> PRIOR FILING DATE: 1999-03-05
24 <150> PRIOR APPLICATION NUMBER: PCT/US98/24855
25 <151> PRIOR FILING DATE: 1998-11-20
27 <150> PRIOR APPLICATION NUMBER: US 60/066,364
28 <151> PRIOR FILING DATE: 1997-11-21
30 <150> PRIOR APPLICATION NUMBER: US 60/078,936
31 <151> PRIOR FILING DATE: 1998-03-20
33 <150> PRIOR APPLICATION NUMBER: PCT/US98/19437
34 <151> PRIOR FILING DATE: 1998-09-17
36 <160> NUMBER OF SEQ ID NOS: 30
38 <170> SOFTWARE: FastSEQ for Windows Version 4.0
40 <210> SEQ ID NO: 1
41 <211> LENGTH: 299
42 <212> TYPE: PRT
43 <213> ORGANISM: Homo sapiens
45 <400> SEQUENCE: 1
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48 Leu Ala Ile Leu Leu Cys Ser Leu Ala Leu Gly Ser Val Thr Val His
49               20                  25                  30
50 Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu
51           35                  40                  45
52 Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe
53       50                  55                  60
54 Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr
55 65                  70                  75                  80
56 Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe
57                   85                  90                  95
58 Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser
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Does Not Comply
Corrected Diskette Needed
(pg.3-4)

RAW SEQUENCE LISTING   DATE: 04/06/2005
PATENT APPLICATION: US/10/785,220A   TIME: 14:13:57

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Output Set: N:\CRF4\04062005\J785220A.raw

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60 Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val
61         115                 120                 125
62 Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr
63         130                 135                 140
64 Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro
65 145                 150                 155                 160
66 Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn
67                 165                 170                 175
68 Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro
69             180                 185                 190
70 Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly
71             195                 200                 205
72 Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser
73         210                 215                 220
74 Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val
75 225                 230                 235                 240
76 Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly
77                 245                 250                 255
78 Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly
79             260                 265                 270
80 Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala Arg Ser Glu
81             275                 280                 285
82 Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val
83         290                 295
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87 <211> LENGTH: 321
88 <212> TYPE: PRT
89 <213> ORGANISM: Homo sapiens
91 <400> SEQUENCE: 2
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93 1               5                   10                  15
94 Thr Tyr Gly Arg Pro Ile Leu Glu Val Pro Glu Ser Val Thr Gly Pro
95             20                  25                  30
96 Trp Lys Gly Asp Val Asn Leu Pro Cys Thr Tyr Asp Pro Leu Gln Gly
97         35                  40                  45
98 Tyr Thr Gln Val Leu Val Lys Trp Leu Val Gln Arg Gly Ser Asp Pro
99     50                  55                  60
100 Val Thr Ile Phe Leu Arg Asp Ser Ser Gly Asp His Ile Gln Gln Ala
101 65                  70                  75                  80
102 Lys Tyr Gln Gly Arg Leu His Val Ser His Lys Val Pro Gly Asp Val
103             85                  90                  95
104 Ser Leu Gln Leu Ser Thr Leu Glu Met Asp Asp Arg Ser His Tyr Thr
105             100                 105                 110
106 Cys Glu Val Thr Trp Gln Thr Pro Asp Gly Asn Gln Val Val Arg Asp
107         115                 120                 125
108 Lys Ile Thr Glu Leu Arg Val Gln Lys Leu Ser Val Ser Lys Pro Thr
109     130                 135                 140
110 Val Thr Thr Gly Ser Gly Tyr Gly Phe Thr Val Pro Gln Gly Met Arg
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/785,220A

DATE: 04/06/2005
TIME: 14:13:57

Input Set : A:\39780-1216.TXT
Output Set: N:\CRF4\04062005\J785220A.raw

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111 145                 150                 155                 160
112 Ile Ser Leu Gln Cys Gln Ala Arg Gly Ser Pro Pro Ile Ser Tyr Ile
113             165                 170                 175
114 Trp Tyr Lys Gln Gln Thr Asn Asn Gln Glu Pro Ile Lys Val Ala Thr
115             180                 185                 190
116 Leu Ser Thr Leu Leu Phe Lys Pro Ala Val Ile Ala Asp Ser Gly Ser
117             195                 200                 205
118 Tyr Phe Cys Thr Ala Lys Gly Gln Val Gly Ser Glu Gln His Ser Asp
119     210                 215                 220
120 Ile Val Lys Phe Val Val Lys Asp Ser Ser Lys Leu Leu Lys Thr Lys
121 225                 230                 235                 240
122 Thr Glu Ala Pro Thr Thr Met Thr Tyr Pro Leu Lys Ala Thr Ser Thr
123             245                 250                 255
124 Val Lys Gln Ser Trp Asp Trp Thr Thr Asp Met Asp Gly Tyr Leu Gly
125             260                 265                 270
126 Glu Thr Ser Ala Gly Pro Gly Lys Ser Leu Pro Val Phe Ala Ile Ile
127             275                 280                 285
128 Leu Ile Ile Ser Leu Cys Cys Met Val Val Phe Thr Met Ala Tyr Ile
129     290                 295                 300
130 Met Leu Cys Arg Lys Thr Ser Gln Gln Glu His Val Tyr Glu Ala Ala
131 305                 310                 315                 320
132 Arg
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137 <211> LENGTH: 390
138 <212> TYPE: DNA
139 <213> ORGANISM: Artificial Sequence
141 <220> FEATURE:
142 <223> OTHER INFORMATION: Artificial sequence
144 <400> SEQUENCE: 3
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146 ttgtatggtc tctgaggaag gcggcaacag ctatggggag gtcaaggtca agctcatcgt 120
147 gcttgtgcct ccatccaagc ctacagttaa catcccctcc tctgccacca ttgggaaccg 180
148 ggcagtgctg acatgctcag aacaagatgg ttccccacct tctgaataca cctggttcaa 240
149 agatgggata gtgatgccta cgaatcccaa aagcacccgt gccttcagca actcttccta 300
150 tgtcctgaat cccacaacag gagagctggt ctttgatccc ctgtcagcct ctgatactgg 360
151 agaatacagc tgtgaggcac ggaatgggta 390
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155 <212> TYPE: DNA
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: Artificial sequence
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164 tgttgtgcct cttcatattg gcgatcctgt tgtgctccct ggcattgggc agtgttacag 180
165 ttgcactctt ctgaacctga agtcagaatt cctgagaata atcctgtgaa gttgtcctgt 240
166 gcctactcgg gcttttcttc tccccgtgtg gagtggaagt ttgaccaagg agacaccacc 300
167 agactcgttt gctataataa caagatcaca gcttcctatg aggaccgggt gaccttcttg 360
```

[Handwritten annotations:] Pls explain source of genetic material. — Invalid Response

[Handwritten annotation near line 155/156:] Pls explain source of genetic material. — Invalid Response

[Handwritten annotation at bottom:] Pls see item # 11 on error summary sheet.

RAW SEQUENCE LISTING   DATE: 04/06/2005
PATENT APPLICATION: US/10/785,220A   TIME: 14:13:57

Input Set : A:\39780-1216.TXT
Output Set: N:\CRF4\04062005\J785220A.raw

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168 ccaactggta tcaccttcaa gtccgtgaca cgggaagaca ctgggacata cacttgtatg 420
169 gtctctgagg aaggcggcaa cagctatggg gaggtcaagg tcaagctcat cgtgcttgtg 480
170 cctccatcca agcctacagt taacatcccc tcctctgcca ccattgggaa ccgggcagtg 540
171 ctgacatgct cagaacaaga tggttcccca ccttctgaat acacctggtt caaagatggg 600
172 atagtgatgc ctacgaatcc caaaagcacc cgtgccttca gcaactcttc ctatgtcctg 660
173 aatcccacaa caggagagct ggtctttgat cccctgtcag cctctgatac tggagaatac 720
174 agctgt                                                         726
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177 <211> LENGTH: 1503
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Artificial sequence
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186 ccaattgagc accctggaga tggatgaccg gagccactac acgtgtgaag tcacctggca 120
187 gactcctgat ggcaaccaag tcgtgagaga taagattact gagctccgtg tccagaaact 180
188 ctctgtctcc aagcccacag tgacaactgg cagcggttat ggcttcacgg tgccccaggg 240
189 aatgaggatt agccttcaat gccagggttc ggggttctcc tcccatcagt tatatttggt 300
190 ataagcaaca gactaataac cagggaaccc atcaaagtag caaccctaag taccttactc 360
191 ttcaagcctg cggtgatagc cgactcaggc tcctatttct gcactgccaa gggccaggtt 420
192 ggctctgagc agcacagcga cattgtgaag tttgtggtca aagactcctc aaagctactc 480
193 aagaccaaga ctgaggcacc tacaaccatg acataccct tgaaagcaac atctacagtg 540
194 aagcagtcct gggactggac cactgacatg gatggctacc ttggagagac cagtgctggg 600
195 ccaggaaaga gcctgcctgt ctttgccatc atcctcatca tctccttgtg ctgtatggtg 660
196 gtttttacca tggcctatat catgctctgt cggaagacat cccaacaaga gcatgtctac 720
197 gaagcagcca gggcacatgc cagagaggcc aacgactctg agaaaccat gagggtggcc 780
198 atcttcgcaa gtggctgctc cagtgatgag ccaacttccc agaatctggg gcaacaacta 840
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200 cgcccgcctg ctggacacag ttcctctgga ttatgagttt ctggccactg agggcaaaag 960
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202 tgccttctgc atggccttct tccctgctac ctctcttcct ggatagccca aagtgtccgc 1080
203 ctaccaacac tggagccgct gggagtcact ggctttgccc tggaatttgc cagatgcatc 1140
204 tcaagtaagc cagctgctgg atttggctct gggcccttct agtatctctg ccgggggctt 1200
205 ctggtactcc tctctaaata ccagagggaa gatgcccata gcactaggac ttggtcatca 1260
206 tgcctacaga cactattcaa ctttggcatc ttgccaccag aagacccgag gggaggctca 1320
207 gctctgccag ctcagaggac cagctatatc caggatcatt tctctttctt cagggccaga 1380
208 cagctttaa ttgaaattgt tatttcacag gccagggttc agttctgctc ctccactata 1440
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210 aaa                                                           1503
212 <210> SEQ ID NO: 6
213 <211> LENGTH: 319
214 <212> TYPE: PRT
215 <213> ORGANISM: Homo sapiens
217 <400> SEQUENCE: 6
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219 1               5                   10                  15
220 Val Thr Val Asp Ala Ile Ser Val Glu Thr Pro Gln Asp Val Leu Arg
221             20                  25                  30
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[Handwritten annotations: "See error explanation, see item #11 on error summary sheet." with circles around "Artificial Sequence" and "Artificial sequence" and "Invalid response" noted. Also: "The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors." and an arrow pointing to "1503"]

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222 Ala Ser Gln Gly Lys Ser Val Thr Leu Pro Cys Thr Tyr His Thr Ser
223         35                  40                  45
224 Thr Ser Ser Arg Glu Gly Leu Ile Gln Trp Asp Lys Leu Leu Leu Thr
225     50                  55                  60
226 His Thr Glu Arg Val Val Ile Trp Pro Phe Ser Asn Lys Asn Tyr Ile
227 65                  70                  75                  80
228 His Gly Glu Leu Tyr Lys Asn Arg Val Ser Ile Ser Asn Asn Ala Glu
229                 85                  90                  95
230 Gln Ser Asp Ala Ser Ile Thr Ile Asp Gln Leu Thr Met Ala Asp Asn
231             100                 105                 110
232 Gly Thr Tyr Glu Cys Ser Val Ser Leu Met Ser Asp Leu Glu Gly Asn
233             115                 120                 125
234 Thr Lys Ser Arg Val Arg Leu Leu Val Leu Val Pro Pro Ser Lys Pro
235     130                 135                 140
236 Glu Cys Gly Ile Glu Gly Glu Thr Ile Ile Gly Asn Asn Ile Gln Leu
237 145                 150                 155                 160
238 Thr Cys Gln Ser Lys Glu Gly Ser Pro Thr Pro Gln Tyr Ser Trp Lys
239             165                 170                 175
240 Arg Tyr Asn Ile Leu Asn Gln Glu Gln Pro Leu Ala Gln Pro Ala Ser
241             180                 185                 190
242 Gly Gln Pro Val Ser Leu Lys Asn Ile Ser Thr Asp Thr Ser Gly Tyr
243         195                 200                 205
244 Tyr Ile Cys Thr Ser Ser Asn Glu Glu Gly Thr Gln Phe Cys Asn Ile
245     210                 215                 220
246 Thr Val Ala Val Arg Ser Pro Ser Met Asn Val Ala Leu Tyr Val Gly
247 225                 230                 235                 240
248 Ile Ala Val Gly Val Val Ala Ala Leu Ile Ile Ile Gly Ile Ile Ile
249             245                 250                 255
250 Tyr Cys Cys Cys Cys Arg Gly Lys Asp Asp Asn Thr Glu Asp Lys Glu
251             260                 265                 270
252 Asp Ala Arg Pro Asn Arg Glu Ala Tyr Glu Glu Pro Pro Glu Gln Leu
253         275                 280                 285
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255     290                 295                 300
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271 ctcagaccct gtcaccatct ttctacgtga ctcttctgga gaccatatcc agcaggcaaa 360
272 gtaccagggc cgcctgcatg tgagccacaa ggttccagga gatgtatccc tccaattgag 420
273 caccctggag atggatgacc ggagccacta cacgtgtgaa gtcacctggc agactcctga 480
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VERIFICATION SUMMARY   DATE: 04/06/2005
PATENT APPLICATION: US/10/785,220A   TIME: 14:13:58

Input Set : A:\39780-1216.TXT
Output Set: N:\CRF4\04062005\J785220A.raw